



8B
lms
4-3-01

SEQUENCE LISTING

<110> Hayden, Michael R.
Brooks-Wilson, Angela R.
Pimstone, Simon N.

<120> METHODS AND REAGENTS FOR MODULATING
CHOLESTEROL LEVELS

<130> 50110/002005

<140> US 09/526,193

<141> 2000-03-15

<150> 60/124,702

<151> 1999-03-15

<150> 60/138,048

<151> 1999-06-08

<150> 60/139,600

<151> 1999-06-17

<150> 60/151,977

<151> 1999-09-01

<160> 287

<170> FastSEQ for Windows Version 4.0

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<211> 2261

<212> PRT

<213> Homo sapiens

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Leu	Phe	Ile	Phe	Leu	Ile	Leu	Ile	Ser	Val	Arg	Leu	Ser	Tyr	Pro	Pro
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Tyr	Glu	Gln	His	Glu	Cys	His	Phe	Pro	Asn	Lys	Ala	Met	Pro	Ser	Ala
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Gly	Thr	Leu	Pro	Trp	Val	Gln	Gly	Ile	Ile	Cys	Asn	Ala	Asn	Asn	Pro
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Cys	Phe	Arg	Tyr	Pro	Thr	Pro	Gly	Glu	Ala	Pro	Gly	Val	Val	Gly	Asn
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Phe	Asn	Lys	Ser	Ile	Val	Ala	Arg	Leu	Phe	Ser	Asp	Ala	Arg	Arg	Leu
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Leu	Leu	Tyr	Ser	Gln	Lys	Asp	Thr	Ser	Met	Lys	Asp	Met	Arg	Lys	Val
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Leu	Arg	Thr	Leu	Gln	Gln	Ile	Lys	Lys	Ser	Ser	Ser	Asn	Leu	Lys	Leu

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Gln Asp Phe Leu Val	Asp Asn Glu Thr Phe	Ser Gly Phe Leu Tyr His		
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Asn Leu Ser Leu Pro	Lys Ser Thr Val Asp	Lys Met Leu Arg Ala Asp		
	165	170		175
Val Ile Leu His Lys	Val Phe Leu Gln Gly	Tyr Gln Leu His Leu Thr		
	180	185		190
Ser Leu Cys Asn Gly	Ser Lys Ser Glu Glu	Met Ile Gln Leu Gly Asp		
	195	200		205
Gln Glu Val Ser Glu	Leu Cys Gly Leu Pro	Arg Glu Lys Leu Ala Ala		
	210	215		220
Ala Glu Arg Val Leu	Arg Ser Asn Met Asp	Ile Leu Lys Pro Ile Leu		
225	230	235		240
Arg Thr Leu Asn Ser	Thr Ser Pro Phe Pro	Ser Lys Glu Leu Ala Glu		
	245	250		255
Ala Thr Lys Thr Leu	Leu His Ser Leu Gly	Thr Leu Ala Gln Glu Leu		
	260	265		270
Phe Ser Met Arg Ser	Trp Ser Asp Met Arg	Gln Glu Val Met Phe Leu		
	275	280		285
Thr Asn Val Asn Ser	Ser Ser Ser Ser Thr	Gln Ile Tyr Gln Ala Val		
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Ser Arg Ile Val Cys	Gly His Pro Glu Gly	Gly Gly Leu Lys Ile Lys		
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Ser Leu Asn Trp Tyr	Glu Asp Asn Asn Tyr	Lys Ala Leu Phe Gly Gly		
	325	330		335
Asn Gly Thr Glu Glu	Asp Ala Glu Thr Phe	Tyr Asp Asn Ser Thr Thr		
	340	345		350
Pro Tyr Cys Asn Asp	Leu Met Lys Asn Leu	Glu Ser Ser Pro Leu Ser		
	355	360		365
Arg Ile Ile Trp Lys	Ala Leu Lys Pro Leu	Leu Val Gly Lys Ile Leu		
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Tyr Thr Pro Asp Thr	Pro Ala Thr Arg Gln	Val Met Ala Glu Val Asn		
385	390	395		400
Lys Thr Phe Gln Glu	Leu Ala Val Phe His	Asp Leu Glu Gly Met Trp		
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Glu Glu Leu Ser Pro	Lys Ile Trp Thr Phe	Met Glu Asn Ser Gln Glu		
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Met Asp Leu Val Arg	Met Leu Leu Asp Ser	Arg Asp Asn Asp His Phe		
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Trp Glu Gln Gln Leu	Asp Gly Leu Asp Trp	Thr Ala Gln Asp Ile Val		
	450	455		460
Ala Phe Leu Ala Lys	His Pro Glu Asp Val	Gln Ser Ser Asn Gly Ser		
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Val Tyr Thr Trp Arg	Glu Ala Phe Asn Glu	Thr Asn Gln Ala Ile Arg		
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Thr Ile Ser Arg Phe	Met Glu Cys Val Asn	Leu Asn Lys Leu Glu Pro		
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Ile Ala Thr Glu Val	Trp Leu Ile Asn Lys	Ser Met Glu Leu Leu Asp		
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Glu Arg Lys Phe Trp	Ala Gly Ile Val Phe	Thr Gly Ile Thr Pro Gly		
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Ser Ile Glu Leu Pro	His His Val Lys Tyr	Lys Ile Arg Met Asp Ile		
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Asp Asn Val Glu Arg	Thr Asn Lys Ile Lys	Asp Gly Tyr Trp Asp Pro		

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Gly	Pro	Arg	Ala	Asp	Pro	Phe	Glu	Asp	Met	Arg	Tyr	Val	Trp	Gly	Gly		
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Thr	Gly	Thr	Glu	Lys	Lys	Thr	Gly	Val	Tyr	Met	Gln	Gln	Met	Pro	Tyr		
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Pro	Cys	Tyr	Val	Asp	Asp	Ile	Phe	Leu	Arg	Val	Met	Ser	Arg	Ser	Met		
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Pro	Leu	Phe	Met	Thr	Leu	Ala	Trp	Ile	Tyr	Ser	Val	Ala	Val	Ile	Ile		
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Lys	Gly	Ile	Val	Tyr	Glu	Lys	Glu	Ala	Arg	Leu	Lys	Glu	Thr	Met	Arg		
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Ile	Met	Gly	Leu	Asp	Asn	Ser	Ile	Leu	Trp	Phe	Ser	Trp	Phe	Ile	Ser		
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Lys	Leu	Gly	Asn	Leu	Leu	Pro	Tyr	Ser	Asp	Pro	Ser	Val	Val	Phe	Val		
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Ile	Tyr	Phe	Thr	Leu	Tyr	Leu	Pro	Tyr	Val	Leu	Cys	Val	Ala	Trp	Gln		
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Val	Cys	Pro	Gln	His	Asn	Val	Leu	Phe	Asp	Met	Leu	Thr	Val	Glu	Glu		
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His	Ile	Trp	Phe	Tyr	Ala	Arg	Leu	Lys	Gly	Leu	Ser	Glu	Lys	His	Val		

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Lys Leu Lys Ser Lys Thr Ser Gln Leu Ser Gly Gly Met Gln Arg Lys		
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Leu Ser Val Ala Leu Ala Phe Val Gly Gly Ser Lys Val Val Ile Leu		1040
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Asp Glu Pro Thr Ala Gly Val Asp Pro Tyr Ser Arg Arg Gly Ile Trp		1055
	1060	1065
Glu Leu Leu Lys Tyr Arg Gln Gly Arg Thr Ile Ile Leu Ser Thr		1070
	1075	1080
His His Met Asp Glu Ala Asp Val Leu Gly Asp Arg Ile Ala Ile Ile		1085
	1090	1095
Ser His Gly Lys Leu Cys Cys Val Gly Ser Ser Leu Phe Leu Lys Asn		1100
1105	1110	1115
Gln Leu Gly Thr Gly Tyr Tyr Leu Thr Leu Val Lys Lys Asp Val Glu		1120
	1125	1130
Ser Ser Leu Ser Ser Cys Arg Asn Ser Ser Ser Thr Val Ser Tyr Leu		1135
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Lys Lys Glu Asp Ser Val Ser Gln Ser Ser Ser Asp Ala Gly Leu Gly		1150
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Ser Asp His Glu Ser Asp Thr Leu Thr Ile Asp Val Ser Ala Ile Ser		1165
	1170	1175
Asn Leu Ile Arg Lys His Val Ser Glu Ala Arg Leu Val Glu Asp Ile		1180
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Gly His Glu Leu Thr Tyr Val Leu Pro Tyr Glu Ala Ala Lys Glu Gly		1200
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Ala Phe Val Glu Leu Phe His Glu Ile Asp Asp Arg Leu Ser Asp Leu		1215
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Gly Ile Ser Ser Tyr Gly Ile Ser Glu Thr Thr Leu Glu Glu Ile Phe		1230
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Leu Lys Val Ala Glu Glu Ser Gly Val Asp Ala Glu Thr Ser Asp Gly		1245
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Thr Leu Pro Ala Arg Arg Asn Arg Arg Ala Phe Gly Asp Lys Gln Ser		1260
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Cys Leu Arg Pro Phe Thr Glu Asp Asp Ala Ala Asp Pro Asn Asp Ser		1280
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Asp Ile Asp Pro Glu Ser Arg Glu Thr Asp Leu Leu Ser Gly Met Asp		1295
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Gly Lys Gly Ser Tyr Gln Val Lys Gly Trp Lys Leu Thr Gln Gln Gln		1310
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Phe Val Ala Leu Leu Trp Lys Arg Leu Leu Ile Ala Arg Arg Ser Arg		1325
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Lys Gly Phe Phe Ala Gln Ile Val Leu Pro Ala Val Phe Val Cys Ile		1340
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Ala Leu Val Phe Ser Leu Ile Val Pro Pro Phe Gly Lys Tyr Pro Ser		1360
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Leu Glu Leu Gln Pro Trp Met Tyr Asn Glu Gln Tyr Thr Phe Val Ser		1375
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Asn Asp Ala Pro Glu Asp Thr Gly Thr Leu Glu Leu Leu Asn Ala Leu		1390
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Thr Lys Asp Pro Gly Phe Gly Thr Arg Cys Met Glu Gly Asn Pro Ile		1405
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Pro Asp Thr Pro Cys Gln Ala Gly Glu Glu Glu Trp Thr Thr Ala Pro		1420

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Val Pro Gln Thr	Ile Met Asp Leu Phe	Gln Asn Gly Asn Trp Thr Met				
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Gln Asn Pro Ser	Pro Ala Cys Gln Cys Ser Ser Asp Lys	Ile Lys Lys				
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Met Leu Pro Val	Cys Pro Pro Gly Ala Gly Gly Leu	Pro Pro Pro Gln				
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Arg Lys Gln Asn	Thr Ala Asp Ile Leu Gln Asp Leu	Thr Gly Arg Asn				
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Ile Ser Asp Tyr	Leu Val Lys Thr Tyr Val Gln Ile	Ile Ala Lys Ser				
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Leu Lys Asn Lys	Ile Trp Val Asn Glu Phe Arg Tyr Gly	Gly Phe Ser				
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Leu Gly Val Ser	Asn Thr Gln Ala Leu Pro Ser Gln	Glu Val Asn				
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Asp Ala Ile Lys	Gln Met Lys Lys His Leu Lys Leu	Ala Lys Asp Ser				
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Ser Ala Asp Arg	Phe Leu Asn Ser Leu Gly Arg Phe	Met Thr Gly Leu				
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Asp Thr Arg Asn	Asn Val Lys Val Trp Phe Asn Asn Lys	Gly Trp His				
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Ala Ile Ser Ser	Phe Leu Asn Val Ile Asn Asn Ala	Ile Leu Arg Ala				
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Asn Leu Gln Lys	Gly Glu Asn Pro Ser His Tyr Gly	Ile Thr Ala Phe				
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Asn His Pro Leu	Asn Leu Thr Lys Gln Gln Leu Ser	Glu Val Ala Leu				
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Met Thr Thr Ser	Val Asp Val Leu Val Ser Ile Cys	Val Ile Phe Ala				
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Met Ser Phe Val	Pro Ala Ser Phe Val Val Phe Leu	Ile Gln Glu Arg				
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Val Ser Lys Ala	Lys His Leu Gln Phe Ile Ser Gly	Val Lys Pro Val				
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Ile Tyr Trp Leu	Ser Asn Phe Val Trp Asp Met Cys	Asn Tyr Val Val				
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Pro Ala Thr Leu	Val Ile Ile Ile Phe Ile Cys Phe	Gln Gln Lys Ser				
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Tyr Val Ser Ser	Thr Asn Leu Pro Val Leu Ala Leu	Leu Leu Leu Leu				
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Tyr Gly Trp Ser	Ile Thr Pro Leu Met Tyr Pro Ala	Ser Phe Val Phe				
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Ile Gly Ile Asn	Gly Ser Val Ala Thr Phe Val Leu	Glu Leu Phe Thr				
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Asp Asn Lys Leu	Asn Asn Ile Asn Asp Ile Leu Lys	Ser Val Phe Leu				
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Asn Gln Ala Met	Ala Asp Ala Leu Glu Arg Phe Gly	Glu Asn Arg Phe				
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Val Ser Pro Leu	Ser Trp Asp Leu Val Gly Arg Asn	Leu Phe Ala Met				
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<211> 7860
<212> DNA
<213> Homo sapiens

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105

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123

B

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124

B

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125

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126

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127

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129

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(30)

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165

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